



#9

## Sequence Listing for.txt

&lt;110&gt; Stuart Newman, et al.

&lt;120&gt; SPLICE CHOICE ANTAGONISTS AS THERAPEUTIC AGENTS

&lt;130&gt; 51230-00601

&lt;140&gt; 09/849,967

&lt;141&gt; 2001-05-08

&lt;160&gt; 4

&lt;210&gt; 1

&lt;211&gt; 1689

&lt;212&gt; DNA

&lt;213&gt; Chicken

&lt;220&gt;

&lt;223&gt; SEQ ID NO:1 Full length cDNA for chicken hnRNP A1

&lt;400&gt; 1

gcgttcac ccctcagccg gcggcggtga gtgcgcagg ccagcgcgg cgtgggaccg 60  
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gagtttagat acccttccaa aatggctgct attaaggaaag agagagaggt ggaagattac 180  
aagagaaaaa ggaagacgat cagcacaggc catgagccta aggagccaga gcagttgaga 240  
aagctgttca ttggagggtct gagcttcgag acgacggatg atagctttag agagcactt 300  
aaaaaatggg gcacactcac ggactgtgtg gtatgagag acccacaaac aaaacgttcc 360  
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cgaccacata aggtggatgg acgtgtggtt gaaccaaaga gagcagttc aaggaggat 480  
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gaagtcatgg aagacagaca aagtggaaag aaaagaggt tcgctttgt aacttttgat 660  
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cgtgggggtg gttcaggcaa cttcatgggt cgtggaaatt ttggaggtgg tggagggaaac 840  
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aaaaaaaaaa 1689

&lt;210&gt; 2

&lt;211&gt; 378

&lt;212&gt; PRT

&lt;213&gt; Chicken

&lt;220&gt;

&lt;223&gt; SEQ ID NO:2 Amino acid sequence of chicken hnRNP A1

&lt;400&gt; 2

Met Ala Ala Ile Lys Glu Glu Arg Glu Val Glu Asp Tyr Lys Arg Lys  
1 5 10 15

Sequence Listing for.txt

Arg Lys Thr Ile Ser Thr Gly His Glu Pro Lys Glu Pro Glu Gln Leu  
 20 25 30  
 Arg Lys Leu Phe Ile Gly Gly Leu Ser Phe Glu Thr Thr Asp Asp Ser  
 35 40 45  
 Leu Arg Glu His Phe Glu Lys Trp Gly Thr Leu Thr Asp Cys Val Val  
 50 55 60  
 Met Arg Asp Pro Gln Thr Lys Arg Ser Arg Gly Phe Gly Phe Val Thr  
 65 70 75 80  
 Tyr Ser Cys Val Glu Glu Val Asp Ala Ala Met Ser Ala Arg Pro His  
 85 90 95  
 Lys Val Asp Gly Arg Val Val Glu Pro Lys Arg Ala Val Ser Arg Glu  
 100 105 110  
 Asp Ser Val Lys Pro Gly Ala His Leu Thr Val Lys Ile Phe Val  
 115 120 125  
 Gly Gly Ile Lys Glu Asp Thr Glu Glu Tyr Asn Leu Arg Gly Tyr Phe  
 130 135 140  
 Glu Thr Tyr Gly Lys Ile Glu Thr Ile Glu Val Met Glu Asp Arg Gln  
 145 150 155 160  
 Ser Gly Lys Lys Arg Gly Phe Ala Phe Val Thr Phe Asp Asp His Asp  
 165 170 175  
 Thr Val Asp Lys Ile Val Val Gln Lys Tyr His Thr Ile Asn Gly His  
 180 185 190  
 Asn Cys Glu Asp Lys Lys Ala Leu Ser Lys Gln Glu Met Gln Thr Ala  
 195 200 205  
 Ser Ser Gln Arg Gly Arg Gly Gly Ser Gly Asn Phe Met Gly Arg  
 210 215 220  
 Gly Asn Phe Gly Gly Gly Asn Phe Gly Arg Gly Gly Asn Phe  
 225 230 235 240  
 Gly Gly Arg Gly Gly Tyr Gly Gly Gly Gly Gly Gly Ser Arg  
 245 250 255  
 Gly Ser Phe Gly Gly Asp Gly Tyr Asn Gly Phe Gly Asp Gly Gly  
 260 265 270  
 Asn Tyr Gly Gly Pro Gly Tyr Gly Ser Arg Gly Gly Tyr Gly Gly  
 275 280 285  
 Gly Gly Gly Pro Gly Tyr Gly Asn Pro Gly Gly Gly Tyr Gly Gly  
 290 295 300  
 Gly Gly Gly Tyr Gly Gly Tyr Asn Glu Gly Gly Asn Phe Gly Gly  
 305 310 315 320  
 Asn Tyr Gly Gly Ser Gly Asn Tyr Asn Asp Phe Gly Asn Tyr Ser Gly  
 325 330 335  
 Gln Gln Gln Ser Asn Tyr Gly Pro Met Lys Gly Gly Gly Ser Phe Gly  
 340 345 350  
 Gly Arg Ser Ser Gly Ser Pro Tyr Gly Gly Gly Tyr Gly Ser Gly Ser  
 355 360 365  
 Gly Ser Gly Gly Tyr Gly Gly Arg Arg Phe  
 370 375

<210> 3  
 <211> 314  
 <212> PRT  
 <213> Human

<220>  
 <223> SEQ ID NO:3 Amino acid sequence of human hnRNP A1

<400> 3  
 Pro Lys Glu Pro Glu Gln Leu Arg Lys Leu Phe Ile Gly Gly Leu Ser  
 1 5 10 15  
 Phe Glu Thr Thr Asp Glu Ser Leu Arg Ser His Phe Glu Gln Trp Gly  
 20 25 30  
 Thr Leu Thr Asp Cys Val Val Met Arg Asp Pro Asn Thr Lys Arg Ser  
 35 40 45

Sequence Listing for.txt

Arg Gly Phe Gly Phe Val Thr Tyr Ala Thr Val Glu Glu Val Asp Ala  
 50 55 60  
 Ala Met Asn Ala Arg Pro His Lys Val Asp Gly Arg Val Val Glu Pro  
 65 70 75 80  
 Lys Arg Ala Val Ser Arg Glu Asp Ser Gln Arg Pro Gly Ala His Leu  
 85 90 95  
 Thr Val Lys Ile Phe Val Gly Gly Ile Lys Glu Asp Thr Glu Glu  
 100 105 110  
 His His Leu Arg Asp Tyr Phe Glu Gln Tyr Gly Lys Ile Glu Val Ile  
 115 120 125  
 Glu Ile Met Thr Asp Arg Gly Ser Gly Lys Lys Arg Gly Phe Ala Phe  
 130 135 140  
 Val Thr Phe Asp Asp His Asp Ser Val Asp Lys Ile Val Ile Gln Lys  
 145 150 155 160  
 Tyr His Thr Val Asn Gly His Asn Cys Glu Val Arg Lys Ala Leu Ser  
 165 170 175  
 Lys Gln Glu Met Ala Ser Ala Ser Ser Gln Arg Gly Arg Ser Gly  
 180 185 190  
 Ser Gly Asn Phe Gly Gly Arg Gly Gly Phe Gly Gly Asn Asp  
 195 200 205  
 Asn Phe Gly Arg Gly Gly Asn Phe Ser Gly Arg Gly Gly Phe Gly Gly  
 210 215 220  
 Ser Arg Gly Gly Gly Tyr Gly Ser Gly Asp Gly Tyr Asn Gly  
 225 230 235 240  
 Phe Gly Asn Asp Gly Ser Asn Phe Gly Gly Ser Tyr Asn Asp  
 245 250 255  
 Phe Gly Asn Tyr Asn Asn Gln Ser Ser Asn Phe Gly Pro Met Lys Gly  
 260 265 270  
 Gly Asn Phe Gly Gly Arg Ser Ser Gly Pro Tyr Gly Gly Gly Gln  
 275 280 285  
 Tyr Phe Ala Lys Pro Arg Asn Gln Gly Gly Tyr Gly Gly Ser Ser Ser  
 290 295 300  
 Ser Ser Ser Tyr Gly Ser Gly Arg Arg Phe  
 305 310

<210> 4  
 <211> 1136  
 <212> DNA  
 <213> Chicken

<220>  
 <223> SEQ ID NO:4 Open reading frame of cDNA for chicken hnRNP A1

<400> 4  
 aatggctgct attaaggaag agagagaggt ggaagattac aagagaaaaa ggaagacgt 60  
 cagcacaggc catgagccta aggagccaga gcagttgaga aagctgttca ttggaggtct 120  
 gagcttcgag acgacggatg atagcttgag agagcactt gaaaaatggg gcacactcac 180  
 ggactgtgtg gtgatgagag acccacaaac aaaacgttcc agaggcttg gctttgttac 240  
 ttactcttgc gtggaagagg tggatgcggc catgagcgct cgaccacata aggtggatgg 300  
 acgtgtggtt gaaccaaaga gaggcatttc aagggaggat tctgtaaagc ctggggcgca 360  
 tctcacagta aagaaaaat ttgttggtgg cattaaagaa gatacagaag aatataattt 420  
 aagggggtac tttgaaacat atggcaagat cgaaacgtata gaagtcatgg aagacagaca 480  
 aagtggaaag aaaagaggct tcgctttgt aactttgtat gatcacgata cagttgataa 540  
 aattgttgtt cagaaatacc atactataaa tggtcataaac tgcaagata aaaaagcact 600  
 ctcaaaacaa gagatgcaga ctgccagctc tcagagaggt cgtgggggtg gttcaggcaa 660  
 cttcatgggt cgtggaaatt ttggaggtgg tggagggaaac tttggccgag gaggaaactt 720  
 tggtggaaaga ggaggctatg ggggtgggtgg tggcgggtgtt gggagcagag gaagctttgg 780  
 gggtgggtat ggatacaacg gatttggtaa tggtggcaac tatggaggtg gtcctggcta 840  
 tggcagcaga ggggtttagt gtgggtgggtgg aggaccagga tatggaaacc caggtggtgg 900  
 atatggaggt ggaggaggag gatatggtgg ctacaatgaa ggaggcaatt ttggaggtgg 960  
 taattatgga ggcagtggaa actacaatga ctttggtaac tacagtggac agcagcagtc 1020  
 caattacggt cccatgaaag gtggtggcag ttttgggtgtt agaagttcag gcagtcctta 1080

Sequence Listing for.txt

tggtgttgt tatggatctg gaagtggaag tgggggctat ggtggtagaa gattct 1136